

SEQUENCE LISTING

<110> Virca, Duke
Bird, Timothy A.
Anderson, Dirk M.
Marken, John S.

<120> Human cDNAs Encoding Polypeptides Having Kinase Functions

<130> 2877-US

<160> 16

<170> PatentIn Ver. 2.0

<210> 1

<211> 181

<212> DNA

<213> Homo sapiens

<400> 1

```
gtacgccatg aaggtgctgc gcaaggcggc gctgggtgcag cgcgccaaga cgcaagagca 60
cacgcgcacc gagcgctcgg tgctggagct ggtgcgccag gcgcccttcc tggtcacgct 120
gcactacgct ttccagacgg atgccaagct gcacctcatc ctggactatg tgagcggcgg 180
g                                                    181
```

<210> 2

<211> 221

<212> DNA

<213> Homo sapiens

<400> 2

```
cccagagagg gccacatcag accgcctccg acttcgtgcg ggactcggcg gccagccacc 60
aggcggagacc cgaggcgtag gagcggcgcg tgtgtctcct gcttctgcaa ctctgcaacg 120
ggctggagca cctgaaggag cacgggatca tccaccggga cctgtgcctg gagaacctgc 180
tgctgggtgca ctgcaccctc caggccggcc ccggggccgc c                                                    221
```

<210> 3

<211> 1085

<212> DNA

<213> Homo sapiens

<400> 3

```
cgggcagggc tggagctggg ctgggatccc gagctcggca gcagcgcagc gggccggccc 60
acctgctggt gccctggagg ctctgagccc cggcggcgcc cgggcccacg cggaacgacg 120
gggcgagatg cgagccaccc ctctggctgc tctgcgggt tccctgtcca ggaagaagcg 180
gttggagttg gatgacaact tagataccga gcgtcccgtc cagaaacgag ctcgaaagtgg 240
gccccagccc agactgcccc cctgcctgtt gcccctgagc ccacctactg ctccagatcg 300
tgcaactgct gtggccactg cctcccgtct tgggcccctat gtctctctgg agcccgagga 360
gggcggggcg gcctaccagg ccctgcactg ccctacaggc actgagtata cctgcaagggt 420
gtaccccgtc caggaagccc tggccgtgct ggagccctac gcgcggctgc ccccgccaca 480
gcatgtggct cggcccactg aggtcctggc tggtagccag ctctctctacg cctttttcac 540
tcggacccat ggggacatgc acagcctggt gogaagccgc caccgtatcc ctgagcctga 600
ggctgcgctg ctcttcggcc agatggccac cgccctggcg cactgtcacc agcacgggtct 660
ggtcctgcgt gatctcaagc tgtgtcgtt tgtcttcgct gaccgtgaga ggaagaagct 720
ggtgctggag aacctggagg actcctgcgt gctgactggg ccagatgatt ccctgtggga 780
caagcacgcg tgcccagcct acgtgggacc tgagatactc agctcacggg cctcatactc 840
gggcaaggca gccgatgtct ggagcctggg cgtggcgctc ttcaccatgc tggccggcca 900
ctaccccttc caggactcgg agcctgtcct gctcttcggc aagatccgcc gcggggccta 960
cgccctgcct gcaggcctct cgcccctgc ccgctgtctg gttcgtctgc tccttcgtcg 1020
ggagccagct gaacggctca cagccacagg catcctctctg cacccttggc tgcgacagga 1080
```

cccgga

1085

<210> 4

<211> 388

<212> DNA

<213> Homo sapiens

<400> 4

cagcgagaag	ccgacatgca	tcgcctcttc	aatcacccca	acatccttcg	cctcgtggct	60
tactgtctga	gggaacgggg	tgctaagcat	gaggcctggc	tgctgctacc	attcttcaag	120
agaggtacgc	tgtggaatga	gatagaaagg	ctgaaggaca	aaggcaactt	cctgaccgag	180
gatcaaattc	tttggctgct	gctggggatc	tgagagggcc	ttgaggccat	tcatgccaaag	240
ggttatgcct	acagagactt	gaagcccacc	aatatattgc	ttggagatga	ggggcagcca	300
gttttaatgg	acttgggttc	catgaatcaa	gcatgcatcc	atgtggaggg	ctcccgccag	360
gctctgacct	tcgaggactg	ggcagccc				388

<210> 5

<211> 1555

<212> DNA

<213> Homo sapiens

<400> 5

atgctaacta	gtttaaacag	atcttgggaac	gagacgacct	gctgtggaag	agcgagcttt	60
ttggaactgt	gcacggggaca	gattggacgc	acacccctcg	ggaggcgcg	agggcatggaa	120
aatttgaagc	atattatcac	ccttggccag	gtcatccaca	aacgggtgtga	agagatgaaa	180
tactgcaaga	aacagtgccg	gcgcctgggc	caccgcgtcc	tcggcctgat	caagcctctg	240
gagatgctcc	aggaccaagg	aaagaggagc	gtgccctctg	agaagttaac	cacagccatg	300
aaccgcttca	aggctgccct	ggaggaggct	aatggggaga	tagaaaagtt	cagcaataga	360
tccaatatct	gcagggtttct	aacagcaagc	caggacaaaa	tactcttcaa	ggacgtgaac	420
aggaagctga	gtgatgtctg	gaaggagctc	tcgctgttac	ttcagggttg	gcaacgcatg	480
cctgtttcac	ccataagcca	aggagcgtcc	tgggcacagg	aagatcagca	ggatgcagac	540
gaagacaggc	gagctttcca	gatgctaaga	agagataatg	aaaaaataga	agcttcactg	600
agacgattag	aaatcaacat	gaaagaaatc	aaggaaactt	tgaggcagta	tttaccacca	660
aatgcatgc	aggagatccc	gcaagagcaa	atcaaggaga	tcaagaagga	gcagctttca	720
ggatccccgt	ggattctgct	aagggaaaaat	gaagtcagca	cactttataa	aggagaatac	780
cacagagctc	cagtggccat	aaaagtattc	aaaaaactcc	aggctggcag	cattgcaata	840
gtgaggcaga	ctttcaataa	ggagatcaaa	accatgaaga	aattcgaatc	tcccaacatc	900
ctgcgtatat	ttgggatttg	cattgatgaa	acagtgactc	cgctcaatt	ctccattgtc	960
atggagtact	gtgaactcgg	gaccctgagg	gagctgttgg	atagggaaaa	agacctcaca	1020
cttggaagc	gcatggctct	agtcctgggg	gcagcccag	gcctataccg	gctacaccat	1080
tcagaagcac	ctgaactcca	cggaaaaatc	agaagctcaa	acttcctggg	aactcaaggc	1140
taccaagtga	agcttgacag	atgtgagttg	aggaaaacac	agacttccat	gagtttggga	1200
actacgagag	aaaagacaga	cagagtcaaa	tctacagcat	atctctcacc	tcaggaactg	1260
gaagatgtat	tttatcaata	tgatgtaaag	tctgaaatat	acagcttttg	aatcgctctc	1320
tgggaaatcg	ccactggaga	tatcccgttt	caaggctgta	attctgagaa	gatccgcaag	1380
ctgggtggctg	tgaagcggca	gcaggagcca	ctgggtgaag	actgcccttc	agagctgagg	1440
gagatcattg	atgagtgcgg	ggcagcaggt	cgtctcgttc	caagatctgt	agcggccgcc	1500
cgggccgtcg	acgttttaac	gcgtggccct	cgagagggtt	tccgatccgg	tcgat	1555

<210> 6

<211> 1498

<212> DNA

<213> Homo sapiens

<400> 6

cttcccgtcg	gacgtggagt	acggaggccc	agaccggagg	tgcccgcctc	cgccctaccc	60
gaagcacctg	ctgctgcgca	gcaagtcgga	gcagtacgac	ctggacagcc	tgtgcgcagg	120
catggagcag	agcctccgtg	cgggccccaa	cgagcccgag	ggcggcgaca	agagccgcaa	180
aagcgccaag	ggggacaaaag	gcggaaaagg	taaaaagcag	attcagacct	ctcccgttcc	240
cgtccgcaaa	aacagcagag	acgaagagaa	gagagagtca	cgcatacaaga	gctactcgcc	300
atacgccttt	aagttcttca	tggagcagca	cgtggagaat	gtcatcaaaa	cctaccagca	360

```

gaagggttaac cggagggtgc agctggagca agaaatggcc aaagctggac tctgtgaagc 420
tgagcaggag cagatgcgga agatcctcta ccagaaagag tctaattaca acaggttaaa 480
gagggccaag atggacaagt ctatgtttgt caagatcaaa accctgggga tcggtgcctt 540
tgagagaagtg tgccttgctt gtaagggtgga cactcacgcc ctgtacgcca tgaagaccct 600
aaggaaaaag gatgtcctga accggaatca ggtggcccac gtcaaggccg agagggacat 660
cctggcccgag gcagacaatg agtgggtggt caaactctac tactccttcc aagacaaaga 720
cagcctgtac tttgtgatgg actacatccc tgggtggggac atgatgagcc tgctgatccg 780
gatggagggtc ttccctgagc acctggcccg gttctacatc gcagagctga ctttggccat 840
tgagagtgtc cacaagatgg gcttcatcca ccgagacatc aagcctgata acattttgat 900
agatctggat ggtcacatta aactcacaga ttctggcctc tgcactgggt tcaggtggac 960
tcacaattcc aaatattacc agaaaggagg ccatgtcaga caggacagca tggagcccag 1020
cgacctctgg gatgatgtgt ctaactgtcg gtgtggggac aggctgaaga ccctagagca 1080
gagggcgcgag aagcagcacc agaggtgcct ggcacattca ctggtgggga ctccaaacta 1140
catcgacccc gaggtgctcc tccgcaaagg gtacactcaa ctctgtgact ggtggagtgt 1200
tgagagtgatt ctcttcgaga tgctgggtgg gcagccgccc tttttggcac ctactcccac 1260
agaaaccagg ctgaagggtga tcaactggga gaacacgctc cacattccag cccaggtgaa 1320
gctgagccct gagggccagg acctcatcac caagctgtgc tgctccgcag accaccgcct 1380
ggggcggaat ggggcccgat acctgaaggc ccaccccttc ttcagcgcca ttgacttctc 1440
cagtgcacatc cggaagcatc cagcccccta cgttcccacc atcagccacc ccatggag 1498

```

<210> 7

<211> 60

<212> PRT

<213> Homo sapiens

<400> 7

```

Tyr Ala Met Lys Val Leu Arg Lys Ala Ala Leu Val Gln Arg Ala Lys
  1              5              10              15

```

```

Thr Gln Glu His Thr Arg Thr Glu Arg Ser Val Leu Glu Leu Val Arg
      20              25              30

```

```

Gln Ala Pro Phe Leu Val Thr Leu His Tyr Ala Phe Gln Thr Asp Ala
      35              40              45

```

```

Lys Leu His Leu Ile Leu Asp Tyr Val Ser Gly Gly
      50              55              60

```

<210> 8

<211> 73

<212> PRT

<213> Homo sapiens

<400> 8

```

Arg Glu Val Pro His Gln Thr Ala Ser Asp Phe Val Arg Asp Ser Ala
  1              5              10              15

```

```

Ala Ser His Gln Ala Glu Pro Glu Ala Tyr Glu Arg Arg Val Cys Phe
      20              25              30

```

```

Leu Leu Leu Gln Leu Cys Asn Gly Leu Glu His Leu Lys Glu His Gly
      35              40              45

```

```

Ile Ile His Arg Asp Leu Cys Leu Glu Asn Leu Leu Leu Val His Cys
      50              55              60

```

```

Thr Leu Gln Ala Gly Pro Gly Pro Ala
      65              70

```

<210> 9
 <211> 360
 <212> PRT
 <213> Homo sapiens

<400> 9

Gly	Gln	Gly	Trp	Ser	Trp	Ala	Gly	Ile	Pro	Ser	Ser	Ala	Ala	Ala	Gln
1				5					10					15	
Arg	Ala	Gly	Pro	Pro	Ala	Gly	Ala	Leu	Glu	Ala	Leu	Ser	Pro	Gly	Gly
			20					25					30		
Ala	Arg	Ala	His	Ala	Glu	Arg	Arg	Gly	Glu	Met	Arg	Ala	Thr	Pro	Leu
		35					40					45			
Ala	Ala	Pro	Ala	Gly	Ser	Leu	Ser	Arg	Lys	Lys	Arg	Leu	Glu	Leu	Asp
	50					55					60				
Asp	Asn	Leu	Asp	Thr	Glu	Arg	Pro	Val	Gln	Lys	Arg	Ala	Arg	Ser	Gly
65					70					75					80
Pro	Gln	Pro	Arg	Leu	Pro	Pro	Cys	Leu	Leu	Pro	Leu	Ser	Pro	Pro	Thr
				85					90						95
Ala	Pro	Asp	Arg	Ala	Thr	Ala	Val	Ala	Thr	Ala	Ser	Arg	Leu	Gly	Pro
			100					105					110		
Tyr	Val	Leu	Leu	Glu	Pro	Glu	Glu	Gly	Gly	Arg	Ala	Tyr	Gln	Ala	Leu
		115					120					125			
His	Cys	Pro	Thr	Gly	Thr	Glu	Tyr	Thr	Cys	Lys	Val	Tyr	Pro	Val	Gln
	130					135					140				
Glu	Ala	Leu	Ala	Val	Leu	Glu	Pro	Tyr	Ala	Arg	Leu	Pro	Pro	His	Lys
145					150					155					160
His	Val	Ala	Arg	Pro	Thr	Glu	Val	Leu	Ala	Gly	Thr	Gln	Leu	Leu	Tyr
			165						170					175	
Ala	Phe	Phe	Thr	Arg	Thr	His	Gly	Asp	Met	His	Ser	Leu	Val	Arg	Ser
			180					185					190		
Arg	His	Arg	Ile	Pro	Glu	Pro	Glu	Ala	Ala	Val	Leu	Phe	Arg	Gln	Met
	195						200					205			
Ala	Thr	Ala	Leu	Ala	His	Cys	His	Gln	His	Gly	Leu	Val	Leu	Arg	Asp
	210					215					220				
Leu	Lys	Leu	Cys	Arg	Phe	Val	Phe	Ala	Asp	Arg	Glu	Arg	Lys	Lys	Leu
225					230					235					240
Val	Leu	Glu	Asn	Leu	Glu	Asp	Ser	Cys	Val	Leu	Thr	Gly	Pro	Asp	Asp
			245						250					255	
Ser	Leu	Trp	Asp	Lys	His	Ala	Cys	Pro	Ala	Tyr	Val	Gly	Pro	Glu	Ile
			260					265					270		
Leu	Ser	Ser	Arg	Ala	Ser	Tyr	Ser	Gly	Lys	Ala	Ala	Asp	Val	Trp	Ser
	275						280					285			
Leu	Gly	Val	Ala	Leu	Phe	Thr	Met	Leu	Ala	Gly	His	Tyr	Pro	Phe	Gln

290

295

300

Asp Ser Glu Pro Val Leu Leu Phe Gly Lys Ile Arg Arg Gly Ala Tyr
 305 310 315 320

Ala Leu Pro Ala Gly Leu Ser Ala Pro Ala Arg Cys Leu Val Arg Cys
 325 330 335

Leu Leu Arg Arg Glu Pro Ala Glu Arg Leu Thr Ala Thr Gly Ile Leu
 340 345 350

Leu His Pro Trp Leu Arg Gln Asp
 355 360

<210> 10

<211> 146

<212> PRT

<213> Homo sapiens

<221> UNSURE

<222> (140)..(140)<223> UNSURE

<400> 10

Gln Arg Glu Ala Asp Met His Arg Leu Phe Asn His Pro Asn Ile Leu
 1 5 10 15

Arg Leu Val Ala Tyr Cys Leu Arg Glu Arg Gly Ala Lys His Glu Ala
 20 25 30

Trp Leu Leu Leu Pro Phe Phe Lys Arg Gly Thr Leu Trp Asn Glu Ile
 35 40 45

Glu Arg Leu Lys Asp Lys Gly Asn Phe Leu Thr Glu Asp Gln Ile Leu
 50 55 60

Trp Leu Leu Leu Gly Ile Cys Arg Gly Leu Glu Ala Ile His Ala Lys
 65 70 75 80

Gly Tyr Ala Tyr Arg Asp Leu Lys Pro Thr Asn Ile Leu Leu Gly Asp
 85 90 95

Glu Gly Gln Pro Val Leu Met Asp Leu Gly Ser Met Asn Gln Ala Cys
 100 105 110

Ile His Val Glu Gly Ser Arg Gln Ala Leu Thr Leu Gln Asp Trp Ala
 115 120 125

Ala Gln Arg Cys Thr Ile Ser Tyr Arg Ala Pro Xaa Leu Phe Ser Val
 130 135 140

Gln Ser
 145

<210> 11

<211> 505

<212> PRT

<213> Homo sapiens

<400> 11

Met Leu Thr Ser Leu Asn Arg Ser Trp Asn Glu Thr Thr Cys Cys Gly
 1 5 10 15
 Arg Ala Ser Phe Leu Glu Leu Cys Thr Gly Gln Ile Gly Arg Thr Pro
 20 25 30
 Leu Gly Arg Arg Glu Gly Met Glu Asn Leu Lys His Ile Ile Thr Leu
 35 40 45
 Gly Gln Val Ile His Lys Arg Cys Glu Glu Met Lys Tyr Cys Lys Lys
 50 55 60
 Gln Cys Arg Arg Leu Gly His Arg Val Leu Gly Leu Ile Lys Pro Leu
 65 70 75 80
 Glu Met Leu Gln Asp Gln Gly Lys Arg Ser Val Pro Ser Glu Lys Leu
 85 90 95
 Thr Thr Ala Met Asn Arg Phe Lys Ala Ala Leu Glu Glu Ala Asn Gly
 100 105 110
 Glu Ile Glu Lys Phe Ser Asn Arg Ser Asn Ile Cys Arg Phe Leu Thr
 115 120 125
 Ala Ser Gln Asp Lys Ile Leu Phe Lys Asp Val Asn Arg Lys Leu Ser
 130 135 140
 Asp Val Trp Lys Glu Leu Ser Leu Leu Leu Gln Val Glu Gln Arg Met
 145 150 155 160
 Pro Val Ser Pro Ile Ser Gln Gly Ala Ser Trp Ala Gln Glu Asp Gln
 165 170 175
 Gln Asp Ala Asp Glu Asp Arg Arg Ala Phe Gln Met Leu Arg Arg Asp
 180 185 190
 Asn Glu Lys Ile Glu Ala Ser Leu Arg Arg Leu Glu Ile Asn Met Lys
 195 200 205
 Glu Ile Lys Glu Thr Leu Arg Gln Tyr Leu Pro Pro Lys Cys Met Gln
 210 215 220
 Glu Ile Pro Gln Glu Gln Ile Lys Glu Ile Lys Lys Glu Gln Leu Ser
 225 230 235 240
 Gly Ser Pro Trp Ile Leu Leu Arg Glu Asn Glu Val Ser Thr Leu Tyr
 245 250 255
 Lys Gly Glu Tyr His Arg Ala Pro Val Ala Ile Lys Val Phe Lys Lys
 260 265 270
 Leu Gln Ala Gly Ser Ile Ala Ile Val Arg Gln Thr Phe Asn Lys Glu
 275 280 285
 Ile Lys Thr Met Lys Lys Phe Glu Ser Pro Asn Ile Leu Arg Ile Phe
 290 295 300
 Gly Ile Cys Ile Asp Glu Thr Val Thr Pro Pro Gln Phe Ser Ile Val
 305 310 315 320
 Met Glu Tyr Cys Glu Leu Gly Thr Leu Arg Glu Leu Leu Asp Arg Glu

325					330					335						
Lys	Asp	Leu	Thr	Leu	Gly	Lys	Arg	Met	Val	Leu	Val	Leu	Gly	Ala	Ala	
340					345					350						
Arg	Gly	Leu	Tyr	Arg	Leu	His	His	Ser	Glu	Ala	Pro	Glu	Leu	His	Gly	
355					360					365						
Lys	Ile	Arg	Ser	Ser	Asn	Phe	Leu	Val	Thr	Gln	Gly	Tyr	Gln	Val	Lys	
370					375					380						
Leu	Ala	Gly	Phe	Glu	Leu	Arg	Lys	Thr	Gln	Thr	Ser	Met	Ser	Leu	Gly	
385					390					395					400	
Thr	Thr	Arg	Glu	Lys	Thr	Asp	Arg	Val	Lys	Ser	Thr	Ala	Tyr	Leu	Ser	
405					410					415						
Pro	Gln	Glu	Leu	Glu	Asp	Val	Phe	Tyr	Gln	Tyr	Asp	Val	Lys	Ser	Glu	
420					425					430						
Ile	Tyr	Ser	Phe	Gly	Ile	Val	Leu	Trp	Glu	Ile	Ala	Thr	Gly	Asp	Ile	
435					440					445						
Pro	Phe	Gln	Gly	Cys	Asn	Ser	Glu	Lys	Ile	Arg	Lys	Leu	Val	Ala	Val	
450					455					460						
Lys	Arg	Gln	Gln	Glu	Pro	Leu	Gly	Glu	Asp	Cys	Pro	Ser	Glu	Leu	Arg	
465					470					475					480	
Glu	Ile	Ile	Asp	Glu	Cys	Arg	Ala	Ala	Gly	Arg	Leu	Val	Pro	Arg	Ser	
485					490					495						
Val	Ala	Ala	Ala	Arg	Ala	Val	Asp	Val								
500					505											

<210> 12
 <211> 499
 <212> PRT
 <213> Homo sapiens

<400> 12
 Phe Pro Leu Asp Val Glu Tyr Gly Gly Pro Asp Arg Arg Cys Pro Pro
 1 5 10 15
 Pro Pro Tyr Pro Lys His Leu Leu Leu Arg Ser Lys Ser Glu Gln Tyr
 20 25 30
 Asp Leu Asp Ser Leu Cys Ala Gly Met Glu Gln Ser Leu Arg Ala Gly
 35 40 45
 Pro Asn Glu Pro Glu Gly Gly Asp Lys Ser Arg Lys Ser Ala Lys Gly
 50 55 60
 Asp Lys Gly Gly Lys Asp Lys Lys Gln Ile Gln Thr Ser Pro Val Pro
 65 70 75 80
 Val Arg Lys Asn Ser Arg Asp Glu Glu Lys Arg Glu Ser Arg Ile Lys
 85 90 95
 Ser Tyr Ser Pro Tyr Ala Phe Lys Phe Phe Met Glu Gln His Val Glu

100					105					110					
Asn	Val	Ile	Lys	Thr	Tyr	Gln	Gln	Lys	Val	Asn	Arg	Arg	Leu	Gln	Leu
		115					120					125			
Glu	Gln	Glu	Met	Ala	Lys	Ala	Gly	Leu	Cys	Glu	Ala	Glu	Gln	Glu	Gln
	130					135					140				
Met	Arg	Lys	Ile	Leu	Tyr	Gln	Lys	Glu	Ser	Asn	Tyr	Asn	Arg	Leu	Lys
145					150					155					160
Arg	Ala	Lys	Met	Asp	Lys	Ser	Met	Phe	Val	Lys	Ile	Lys	Thr	Leu	Gly
			165						170					175	
Ile	Gly	Ala	Phe	Gly	Glu	Val	Cys	Leu	Ala	Cys	Lys	Val	Asp	Thr	His
		180						185					190		
Ala	Leu	Tyr	Ala	Met	Lys	Thr	Leu	Arg	Lys	Lys	Asp	Val	Leu	Asn	Arg
		195					200					205			
Asn	Gln	Val	Ala	His	Val	Lys	Ala	Glu	Arg	Asp	Ile	Leu	Ala	Glu	Ala
	210					215					220				
Asp	Asn	Glu	Trp	Val	Val	Lys	Leu	Tyr	Tyr	Ser	Phe	Gln	Asp	Lys	Asp
225					230					235					240
Ser	Leu	Tyr	Phe	Val	Met	Asp	Tyr	Ile	Pro	Gly	Gly	Asp	Met	Met	Ser
			245						250					255	
Leu	Leu	Ile	Arg	Met	Glu	Val	Phe	Pro	Glu	His	Leu	Ala	Arg	Phe	Tyr
			260					265					270		
Ile	Ala	Glu	Leu	Thr	Leu	Ala	Ile	Glu	Ser	Val	His	Lys	Met	Gly	Phe
	275					280						285			
Ile	His	Arg	Asp	Ile	Lys	Pro	Asp	Asn	Ile	Leu	Ile	Asp	Leu	Asp	Gly
	290					295					300				
His	Ile	Lys	Leu	Thr	Asp	Phe	Gly	Leu	Cys	Thr	Gly	Phe	Arg	Trp	Thr
305					310					315					320
His	Asn	Ser	Lys	Tyr	Tyr	Gln	Lys	Gly	Ser	His	Val	Arg	Gln	Asp	Ser
			325						330					335	
Met	Glu	Pro	Ser	Asp	Leu	Trp	Asp	Asp	Val	Ser	Asn	Cys	Arg	Cys	Gly
			340					345					350		
Asp	Arg	Leu	Lys	Thr	Leu	Glu	Gln	Arg	Ala	Arg	Lys	Gln	His	Gln	Arg
		355					360					365			
Cys	Leu	Ala	His	Ser	Leu	Val	Gly	Thr	Pro	Asn	Tyr	Ile	Ala	Pro	Glu
	370					375					380				
Val	Leu	Leu	Arg	Lys	Gly	Tyr	Thr	Gln	Leu	Cys	Asp	Trp	Trp	Ser	Val
385					390					395					400
Gly	Val	Ile	Leu	Phe	Glu	Met	Leu	Val	Gly	Gln	Pro	Pro	Phe	Leu	Ala
			405						410					415	
Pro	Thr	Pro	Thr	Glu	Thr	Gln	Leu	Lys	Val	Ile	Asn	Trp	Glu	Asn	Thr
			420					425					430		

Leu His Ile Pro Ala Gln Val Lys Leu Ser Pro Glu Ala Arg Asp Leu
 435 440 445
 Ile Thr Lys Leu Cys Cys Ser Ala Asp His Arg Leu Gly Arg Asn Gly
 450 455 460
 Ala Asp Asp Leu Lys Ala His Pro Phe Phe Ser Ala Ile Asp Phe Ser
 465 470 475 480
 Ser Asp Ile Arg Lys His Pro Ala Pro Tyr Val Pro Thr Ile Ser His
 485 490 495
 Pro Met Glu

<210> 13
 <211> 375
 <212> DNA
 <213> Homo sapiens

<400> 13
 cttgcaggat ttgagttgag gaaaacacag acttccatga gtttggggaac tacgagagaa 60
 aagacagaca gagtcaaadc tacagcatat ctctcacctc aggaactgga agatgtatatt 120
 tatcaatatg atgtaaagtc tgaaatatac agctttggaa tcgtcctctg ggaaatcgcc 180
 actggagata tcccgtttca aggctgtaat tctgagaaga tccgcaagct ggtggctgtg 240
 aagcggcagc aggagccact ggggtgaagac tgcccttcag agctgcggga gatcattgat 300
 gagtgccggg cccatgatcc ctctgtgcgg ccctctgtgg atgaaatctt aaagaaactc 360
 tccacctttt ctaag 375

<210> 14
 <211> 125
 <212> PRT
 <213> Homo sapiens

<400> 14
 Leu Ala Gly Phe Glu Leu Arg Lys Thr Gln Thr Ser Met Ser Leu Gly
 1 5 10 15
 Thr Thr Arg Glu Lys Thr Asp Arg Val Lys Ser Thr Ala Tyr Leu Ser
 20 25 30
 Pro Gln Glu Leu Glu Asp Val Phe Tyr Gln Tyr Asp Val Lys Ser Glu
 35 40 45
 Ile Tyr Ser Phe Gly Ile Val Leu Trp Glu Ile Ala Thr Gly Asp Ile
 50 55 60
 Pro Phe Gln Gly Cys Asn Ser Glu Lys Ile Arg Lys Leu Val Ala Val
 65 70 75 80
 Lys Arg Gln Gln Glu Pro Leu Gly Glu Asp Cys Pro Ser Glu Leu Arg
 85 90 95
 Glu Ile Ile Asp Glu Cys Arg Ala His Asp Pro Ser Val Arg Pro Ser
 100 105 110
 Val Asp Glu Ile Leu Lys Lys Leu Ser Thr Phe Ser Lys
 115 120 125

<210> 15
 <211> 1961
 <212> DNA
 <213> Homo sapiens

<400> 15
 tccccgctgga cgtggagtag ggaggcccag accggagggtg cccgcctccg ccctaccga 60
 agcacctgct gctgcgcagc aagtcggagc agtacgacct ggacagcctg tgcgcaggca 120
 tggagcagag cctccgtgcg ggcccacacg agcccagggg cggcgacaag agccgcaaaa 180
 gcgccaaggg ggacaaaggc ggaaaggata aaaagcagat tcagacctct cccgttcccc 240
 tccgcaaaaa cagcagagac gaagagaaga gagagtcacg catcaagagc tactcgccat 300
 acgcctttta gttcttcatg gagcagcacg tggagaatgt catcaaaacc taccagcaga 360
 aggttaaccg gaggtctgcag ctggagcaag aaatggccaa agctggactc tgtgaagctg 420
 agcaggagca gatgcggaag atcctctacc agaaagagtc taattacaac aggttaaaga 480
 gggccaagat ggacaagtct atgtttgtca agatcaaaac cctggggatc ggtgcctttg 540
 gagaagtgtg ccttgcttgt aaggtggaca ctcacgccct gtacgccatg aagaccctaa 600
 ggaaaaagga tgtcctgaac cggaaatcagg tggcccacgt caaggccgag agggacatcc 660
 tggccgaggg agacaatgag tgggtgggtc aactctacta ctccctccaa gacaaagaca 720
 gcctgtactt tgtgatggac tacatccctg gtggggacat gatgagcctg ctgatccgga 780
 tggaggtctt ccctgagcac ctggcccggg tctacatcgc agagctgact ttggccattg 840
 agagtgtcca caagatgggc ttcattccacc gagacatcaa gctgataaac attttgatag 900
 atctggatgg tcacattaaa ctcacagatt tcggcctctg cactgggttc aggtggactc 960
 acaattccaa atattaccag aaagggagcc atgtcagaca ggacagcatg gagcccagcg 1020
 acctctggga tgatgtgtct aactgtcggg gtggggacag gctgaagacc ctagagcaga 1080
 gggcgcgga gacagaccag aggtgcctgg cacattcact ggtggggact ccaaactaca 1140
 tcgcacccga ggtgtctctc cgcaaagggt acactcaact ctgtgactgg tggagtgttg 1200
 gagtgattct cttcgagatg ctggtggggc agccgccctt tttggcacct actcccacag 1260
 aaaccagct gaaggtgac aactgggaga acacgctcca cattccagcc caggtgaagc 1320
 tgagccctga ggccaggac ctcattacca agctgtgctg ctccgcagac caccgcctgg 1380
 ggcggaatgg ggccgatgac ctgaaggccc accccttctt cagcgccatt gacttctcca 1440
 gtgacatccg gaagcatcca gcccctacg tttccaccat cagccacccc atggacacct 1500
 cgaatttcga ccccgtagat gaagaaagcc cttggaacga tgccagcgaa ggtagcacca 1560
 aggcctggga cacactcacc tcgcccata acaagcatcc tgagcacgca ttttacgaat 1620
 tcaccttccg aaggttcttt gatgacaatg gctaccctt tcgatgcca aagccttcag 1680
 gagcagaagc ttcacaggct gagagctcag atttagaaag ctctgatctg gtggatcaga 1740
 ctgaaggctg ccagcctgtg tacgtgtaga tgggggcccag gcacccccac cactcgctgc 1800
 ctcccaggtc aggttcccgg agccggtgcc ctcacaggcc aataggggaag ccgagggctg 1860
 ttttgtttta aattagtccg tcgattactt cacttgaaat tctgctcttc accaagaaaa 1920
 cccaaacagg acacttttga aaacagcggg gccgcgaatt c 1961

<210> 16
 <211> 588
 <212> PRT
 <213> Homo sapiens

<400> 16
 Pro Leu Asp Val Glu Tyr Gly Gly Pro Asp Arg Arg Cys Pro Pro Pro
 1 5 10 15
 Pro Tyr Pro Lys His Leu Leu Leu Arg Ser Lys Ser Glu Gln Tyr Asp
 20 25 30
 Leu Asp Ser Leu Cys Ala Gly Met Glu Gln Ser Leu Arg Ala Gly Pro
 35 40 45
 Asn Glu Pro Glu Gly Gly Asp Lys Ser Arg Lys Ser Ala Lys Gly Asp
 50 55 60
 Lys Gly Gly Lys Asp Lys Lys Gln Ile Gln Thr Ser Pro Val Pro Val

65	70					75					80				
Arg	Lys	Asn	Ser	Arg	Asp	Glu	Glu	Lys	Arg	Glu	Ser	Arg	Ile	Lys	Ser
				85					90					95	
Tyr	Ser	Pro	Tyr	Ala	Phe	Lys	Phe	Phe	Met	Glu	Gln	His	Val	Glu	Asn
			100					105					110		
Val	Ile	Lys	Thr	Tyr	Gln	Gln	Lys	Val	Asn	Arg	Arg	Leu	Gln	Leu	Glu
		115					120					125			
Gln	Glu	Met	Ala	Lys	Ala	Gly	Leu	Cys	Glu	Ala	Glu	Gln	Glu	Gln	Met
	130					135					140				
Arg	Lys	Ile	Leu	Tyr	Gln	Lys	Glu	Ser	Asn	Tyr	Asn	Arg	Leu	Lys	Arg
145					150				155						160
Ala	Lys	Met	Asp	Lys	Ser	Met	Phe	Val	Lys	Ile	Lys	Thr	Leu	Gly	Ile
			165						170					175	
Gly	Ala	Phe	Gly	Glu	Val	Cys	Leu	Ala	Cys	Lys	Val	Asp	Thr	His	Ala
			180					185					190		
Leu	Tyr	Ala	Met	Lys	Thr	Leu	Arg	Lys	Lys	Asp	Val	Leu	Asn	Arg	Asn
		195					200					205			
Gln	Val	Ala	His	Val	Lys	Ala	Glu	Arg	Asp	Ile	Leu	Ala	Glu	Ala	Asp
	210					215					220				
Asn	Glu	Trp	Val	Val	Lys	Leu	Tyr	Tyr	Ser	Phe	Gln	Asp	Lys	Asp	Ser
225					230					235					240
Leu	Tyr	Phe	Val	Met	Asp	Tyr	Ile	Pro	Gly	Gly	Asp	Met	Met	Ser	Leu
			245						250					255	
Leu	Ile	Arg	Met	Glu	Val	Phe	Pro	Glu	His	Leu	Ala	Arg	Phe	Tyr	Ile
			260					265					270		
Ala	Glu	Leu	Thr	Leu	Ala	Ile	Glu	Ser	Val	His	Lys	Met	Gly	Phe	Ile
		275					280					285			
His	Arg	Asp	Ile	Lys	Pro	Asp	Asn	Ile	Leu	Ile	Asp	Leu	Asp	Gly	His
	290					295					300				
Ile	Lys	Leu	Thr	Asp	Phe	Gly	Leu	Cys	Thr	Gly	Phe	Arg	Trp	Thr	His
305					310					315					320
Asn	Ser	Lys	Tyr	Tyr	Gln	Lys	Gly	Ser	His	Val	Arg	Gln	Asp	Ser	Met
			325					330						335	
Glu	Pro	Ser	Asp	Leu	Trp	Asp	Asp	Val	Ser	Asn	Cys	Arg	Cys	Gly	Asp
			340					345					350		
Arg	Leu	Lys	Thr	Leu	Glu	Gln	Arg	Ala	Arg	Lys	Gln	His	Gln	Arg	Cys
		355					360					365			
Leu	Ala	His	Ser	Leu	Val	Gly	Thr	Pro	Asn	Tyr	Ile	Ala	Pro	Glu	Val
	370					375					380				
Leu	Leu	Arg	Lys	Gly	Tyr	Thr	Gln	Leu	Cys	Asp	Trp	Trp	Ser	Val	Gly
385					390					395					400

Val	Ile	Leu	Phe	Glu	Met	Leu	Val	Gly	Gln	Pro	Pro	Phe	Leu	Ala	Pro	405	410	415
Thr	Pro	Thr	Glu	Thr	Gln	Leu	Lys	Val	Ile	Asn	Trp	Glu	Asn	Thr	Leu	420	425	430
His	Ile	Pro	Ala	Gln	Val	Lys	Leu	Ser	Pro	Glu	Ala	Arg	Asp	Leu	Ile	435	440	445
Thr	Lys	Leu	Cys	Cys	Ser	Ala	Asp	His	Arg	Leu	Gly	Arg	Asn	Gly	Ala	450	455	460
Asp	Asp	Leu	Lys	Ala	His	Pro	Phe	Phe	Ser	Ala	Ile	Asp	Phe	Ser	Ser	465	470	475
Asp	Ile	Arg	Lys	His	Pro	Ala	Pro	Tyr	Val	Pro	Thr	Ile	Ser	His	Pro	485	490	495
Met	Asp	Thr	Ser	Asn	Phe	Asp	Pro	Val	Asp	Glu	Glu	Ser	Pro	Trp	Asn	500	505	510
Asp	Ala	Ser	Glu	Gly	Ser	Thr	Lys	Ala	Trp	Asp	Thr	Leu	Thr	Ser	Pro	515	520	525
Asn	Asn	Lys	His	Pro	Glu	His	Ala	Phe	Tyr	Glu	Phe	Thr	Phe	Arg	Arg	530	535	540
Phe	Phe	Asp	Asp	Asn	Gly	Tyr	Pro	Phe	Arg	Cys	Pro	Lys	Pro	Ser	Gly	545	550	555
Ala	Glu	Ala	Ser	Gln	Ala	Glu	Ser	Ser	Asp	Leu	Glu	Ser	Ser	Asp	Leu	565	570	575
Val	Asp	Gln	Thr	Glu	Gly	Cys	Gln	Pro	Val	Tyr	Val					580	585	